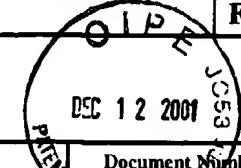


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	Applicant(s): Eric T. BALDWIN et al.	Confirmation No.: 7868
	Filing Date: 29 June 2001	Group: 2621

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Examiner Initial	Document Number	Date	Name	Class	Subclass	Filing Date If Appropriate
	NONE					

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						Yes	No
NE	0 786 519 A2, A3	07/30/98	EPO				
NE	WO 99/47639	09/23/99	WIPO			RECEIVED	
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NE	Adams, "On the Release of the Formyl Group from Nascent Protein," <i>Journal of Molecular Biology</i> , 33(3):571-589 (1968).
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<i>NO</i>	<i>RECEIVED DEC 12 2001 JL53 PATENT &amp; TRADEMARK OFFICE</i>	Blundell, "Metalloproteinase superfamilies and drug design," <i>Nature Structural Biology</i> , 1(2):73-75 (1994).
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N.E. <i>Nashed</i>	5/13/04

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	<b>Filing Date:</b> 29 June 2001	<b>Group:</b> 2621

Examiner Initial	Document Description
<i>Nel</i> 	Martin, "3D Database Searching in Drug Design," <i>Journal of Medicinal Chemistry</i> , 35(12):2145-2154 (1992).
<i>Nel</i> 	Matthews, "Structural Basis of the Action of Thermolysin and Related Zinc Peptidases," <i>Accounts of Chemical Research</i> , 21:333-340 (1988).
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<i>Nel</i> Technology Center 2600	Meinnel et al., "Enzymatic Properties of <i>Escherichia coli</i> Peptide Deformylase," <i>Journal of Bacteriology</i> , 177(7):1883-1887 (1995).
<i>Nel</i> DEC 14 2001	Meng et al., "Automated Docking with Grid-Based Energy Evaluation," <i>Journal of Computational Chemistry</i> , 13(4):505-524 (1992).
<i>Nel</i> 2600	Merritt et al., "Raster3D Version 2.0. A Program for Photorealistic Molecular Graphics," <i>Acta Crystallographica</i> , D50(6):869-873 (1994).
<i>Nel</i>	Miranker et al., "Functionality Maps of Binding Sites: A Multiple Copy Simultaneous Search Method," <i>Proteins: Structure, Function, and Genetics</i> , 11(1):29-34 (1991).
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<i>Nel</i>	Rajagopalan et al., "Peptide Deformylase: A New Type of Mononuclear Iron Protein," <i>Journal of the American Chemical Society</i> , 119(50):12418-12419 (1997).
<i>Nel</i>	Rossmann, ed., <i>The Molecular Replacement Method A Collection of Papers on the Use of Non-Crystallographic Symmetry</i> , Gordon & Breach, Science Publishers, Inc., New York, Title page, publication page, and table of contents only, 6 pages (1972).

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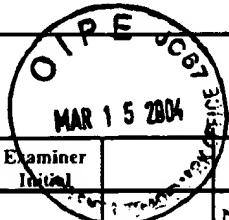
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Examiner Initial	Document Description
<i>ME</i> <i>OIP E</i> <i>REC 12/2001</i> <i>PATENT &amp; TRADEMARK OFFICE</i>	Sack, "CHAIN - A Crystallographic Modeling Program," <i>Journal of Molecular Graphics</i> , 6(4):224-225 (1988).
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<i>ME</i>	Schmitt et al., "Molecular recognition governing the initiation of translation in <i>Escherichia coli</i> . A review," <i>Biochimie</i> , 78(7):543-554 (1996).
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<i>ME</i>	Sheldrick et al., "Structure Solution by Iterative Peaklist Optimization and Tangent Expansion in Space Group P1," <i>Acta Crystallographica, B51</i> (4):423-431 (1995).
<i>ME</i> Technology Center 2800 RECEIVED DEC 1 4 2001	Solbiati et al., "Processing of the N Termini of Nascent Polypeptide Chains Requires Deformylation Prior to Methionine Removal," <i>Journal of Molecular Biology</i> , 290(3):607-614 (1999).
<i>ME</i>	Tatusova et al., "BLAST 2 Sequences, a new tool for comparing protein and nucleotide sequences," <i>FEMS Microbiology Letters</i> , 174(2):247-250 (1999).
<i>ME</i>	Travis, "Proteins and Organic Solvents Make an Eye-Opening Mix," <i>Science</i> , 262(5138):1374 (1993).
<i>ME</i>	Van Duyne et al., "Atomic Structures of the Human Immunophilin FKBP-12 Complexes with FK506 and Rapamycin," <i>Journal of Molecular Biology</i> , 229(1):105-124 (1993).
<i>ME</i>	Weiner et al., "An All Atom Force Field for Simulations of Proteins and Nucleic Acids," <i>Journal of Computational Chemistry</i> , 7(2):230-252 (1986).
<i>ME</i>	Wyckoff et al., eds., "Diffraction Methods for Biological Macromolecules, Part A," <i>Methods in Enzymology</i> , volume 114, Title page, publication page and table of contents only, 3 pages (1985).
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<b>EXAMINER</b> <i>ME</i>	<b>Date Considered</b> <i>5/13/04</i>
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	<b>Application Filing Date:</b> June 29, 2001	<b>Group:</b> 2621
	<b>Information Disclosure Statement mailed:</b>	

**U.S. PATENT DOCUMENTS**

Examiner Initial	Document Number	Date	Name	Class	Subclass	Filing Date If Appropriate
	NONE					

**FOREIGN PATENT DOCUMENTS**

Examiner Initial	Document Number	Date	Country	Class	Subclass	Translation	
						Yes	No
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NC	Chen et al., "Actinonin, a naturally occurring antibacterial agent, is a potent deformylase inhibitor," <i>Biochemistry</i> . 2000;39(6):1256-62.
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<i>Nashad</i>	5/11/04

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EXHIBIT A